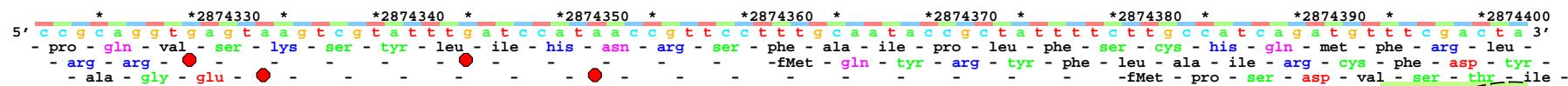
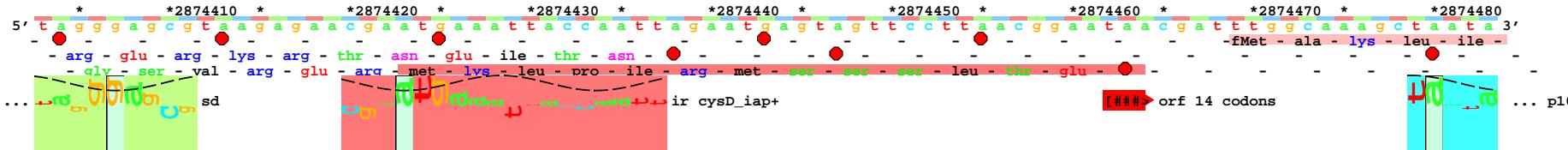


- 1 -

piece 1, NC_000913, cysD_iap+, config: linear, direction: +, begin: 2874322, end: 28746222

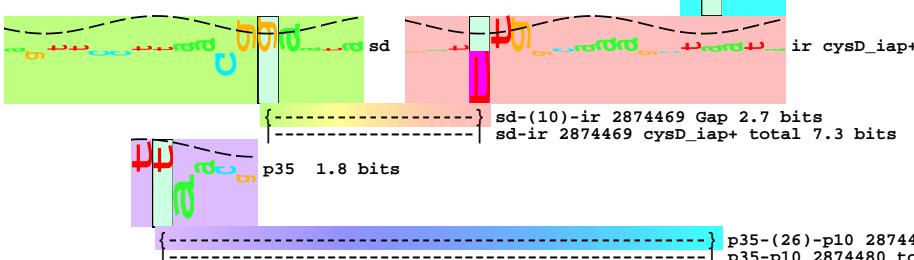


13.cysD



{-----} sd-(16)-ir 2874423 Gap 6.4 bits

|-----| sd-ir 2874423 cysD_iap+ total 7.2 bits



This figure displays a genomic sequence from position *2874490 to *2874560. The sequence is color-coded by codon: A (green), T (blue), C (red), G (yellow). Stop codons are marked with asterisks (*). The first gene (2874490) encodes a protein starting with fMet and ending at met. The second gene (2874500) encodes a protein starting with lys and ending at met. The third gene (2874510) encodes a protein starting with fMet and ending at met. The fourth gene (2874520) encodes a protein starting with fMet and ending at met. The fifth gene (2874530) encodes a protein starting with fMet and ending at met. The sixth gene (2874540) encodes a protein starting with fMet and ending at met. The seventh gene (2874550) encodes a protein starting with fMet and ending at met. The eighth gene (2874560) encodes a protein starting with fMet and ending at met.

The diagram illustrates four sequence segments arranged horizontally:

- p35-(23)-p10**: Total length 5.5 bits, with a gap of 1.4 bits between p35 and p10.
- p35**: Length 4.8 bits.
- p10**: Length 2.4 bits.
- p35-(26)-p10**: Total length 5.2 bits, with a gap of 3.7 bits between p35 and p10.
- p35**: Length 3.8 bits.
- p10**: Length 4.4 bits.

The diagram illustrates two protein domains. The top domain, labeled p35-(23)-p10, has a gap of 1.4 bits and a total length of 7.8 bits. The bottom domain, labeled p35-(22)-p10, has a gap of 2.3 bits and a total length of 7.1 bits. Both domains are shown as blue rectangles with green and red segments indicating specific regions.

